

**CSE- 4029 LAB Assignment - 1&2****Academic year:** 2021-2022**Semester:** WIN**Faculty Name:** Prof. BKSP Kumar raju Alluri**Date:** 18/3/2022**Student name:** M.Taran**Reg. no.:** 19BCE7346**Exploratory Data analysis**

**##Step-1 :** Took Dataset with both categorical and continuous data values and drew graphs Accordingly.

**#link for dataset :** <https://archive.ics.uci.edu/ml/datasets/wholesale+customers>

**#Importing Data**

customers <-

read.csv("D:/users/lenovo/OneDrive/Desktop/19BCE7346/customer\_data.csv",  
stringsAsFactors = FALSE)

head(customers, n = 6)

glimpse(customers)

dim(customers)

**#Understanding data**

ggplot(customers, aes(y = Income)) + geom\_boxplot()

**# Boxplot of the Year of birth variable**

ggplot(customers, aes(Year\_Birth)) + geom\_boxplot()

**#Step-2: Apply any ML- Algorithm without pre-processing**

**#Step-3: Perform Various Categorical and Continuous Encodings**

**#Encoding the categorical features & continuous**

```
# Encoding the categorical features to numeric
customers_copy <- customers_copy %>% mutate(Education =
case_when(Education == "graduate" ~ 1,
           Education == "non-graduate" ~ 0))
customers_copy <- customers_copy %>% mutate(Marital_Status =
case_when(Marital_Status == "Taken" ~ 1,
           Marital_Status == "Single" ~ 0))

str(customers_copy$Education)
str(customers_copy$Marital_Status)

library(caret)
#preprocessing the data
customers_copy_pre <- preProcess(customers_copy[,c(3, 6:17, 25:26)], method =
c("center", "scale"))

#normalizing
customers_copy <- predict(customers_copy_pre, customers_copy[, c(3, 6:17,
25:26)])
summary(customers_copy)

##Step-4: Feature Generation
# Going to drop the rows that have missing income values.
customers <- na.omit(customers)
dim(customers)

library(cluster)
sil_width <- map_dbl(2:10, function(k){
  model <- pam(customers_copy, k = k)
  model$silinfo$avg.width
})
sil_df <- data.frame(
  k = 2:10,
  sil_width = sil_width
)
head(sil_df)
```

```
ggplot(sil_df, aes(k, sil_width)) + geom_line() + scale_x_continuous(breaks = 2:10) +  
labs(y = "Avg sil width")  
trainingSet_scaled <- as.data.frame(scale(trainingSet[,  
getIndependentNumbersOfCol()])))  
testSet_scaled <- as.data.frame(scale(testSet[, getIndependentNumbersOfCol()])))  
dataSet_scaled <- as.data.frame(scale(dataSet[, getIndependentNumbersOfCol()])))
```

### **#Step-5: Dimensionality Reduction**

```
#Using popular method of dimensionality reduction is the principal component  
analysis(PCA)  
library(FactoMineR)
```

```
#Running a PCA.
```

```
customers_copy_pca <- PCA(customers_copy, graph = FALSE)
```

```
#Exploring PCA()
```

```
# Getting the summary of the pca  
summary(customers_copy_pca)
```

```
#Getting the variance of the first 7 new dimensions  
customers_copy_pca$eig[,2][1:7]
```

```
#Getting the cumulative variance  
customers_copy_pca$eig[,3][1:7]
```

```
#Getting the most correlated variables  
dimdesc(customers_copy_pca, axes = 1:2)
```

```
#Tracing variable contributions in customers_pca  
customers_copy_pca$var$contrib
```

```
#Visualising PCA  
library(factoextra)  
#Barplotting the contributions of variables
```

```
fviz_contrib(customers_copy_pca, choice = "var", axes = 1, top = 5)
```

```
#Biplots
```

```
fviz_pca_biplot(customers_copy_pca)
```

### **#Step-6: Creating Missing values and identifying the best missing value technique**

```
#counting the total number of missing values in the data
```

```
library(naniar)
```

```
n_miss(customers)
```

```
# Summarizing missingness in each variable
```

```
miss_var_summary(customers)
```

```
#Pre-Processing Data
```

```
# Going to drop the rows that have missing income values.
```

```
customers <- na.omit(customers)
```

```
dim(customers)
```

```
#Parsing the Dt_Customer as Date object
```

```
#But first i need to make sure that the date is according to ISO 8601 standards  
before converting it to a Date object thats where the function dmy() from the  
lubridate package comes in.
```

```
library(lubridate)
```

```
customers <- customers %>% mutate(Dt_Customer = as.Date(dmy(Dt_Customer)))
```

```
str(customers$Dt_Customer)
```

```
# Dates of the oldest and newest recorded customer
```

```
paste0("The oldest enrolment date of a customer dates to: ",
```

```
min(customers$Dt_Customer))
```

```
paste0("The newest enrolment date of a customer dates to: ",
```

```
max(customers$Dt_Customer))
```

```
#creating a new variable Age from Year of Birth
```

```
customers <- customers %>% mutate(Age = 2021 - Year_Birth)
```

```
customers %>% select(Age) %>% arrange(desc(Age)) %>% top_n(3)
# Max Age is > 100
#Dropping outliers by setting a cap on Income and Age
customers <- customers %>% filter(Income < 600000 & Age < 90)
dim(customers)
```

**#Step-7: Applying k-means algorithm on the final dataset and comparing the performance in step-2 and current performance.**

```
library(corrplot)
```

```
#Getting correlation matrix
cust_cor <- cor(customers[,3:17])
corrplot(cust_cor, method = "color", addCoef.col = "white")
```

```
#The elbow method
```

```
library(purrr)
tot_withinss <- map_dbl(1:10, function(k){
  model <- kmeans(x = customers_copy, centers = k)
  model$tot.withinss
})
```

```
elbow_df <- data.frame(
  k = 1:10,
  tot_withinss = tot_withinss)
head(elbow_df)
#plotting the elbow plot
ggplot(elbow_df, aes(k, tot_withinss)) + geom_line() + scale_x_continuous(breaks =
1:10)
```

```
#Visualizing the top 5 features in the contribution
```

```
#visualizing wines
```

```
customers %>% ggplot(aes(wines)) + geom_histogram(color = "black", fill =
"lightblue") + facet_wrap(vars(cluster))
```

```
#visualizing Income variable
```

```
customers %>% ggplot(aes(Income)) + geom_histogram(color = "black", fill =  
"lightgreen") + facet_wrap(vars(cluster)) +  
geom_vline(aes(xintercept=mean(Income)),color="blue", linetype="dashed", size =  
1)  
#visualizing Total_spent  
customers %>% ggplot(aes(Total_spent)) + geom_histogram(color = "black", fill =  
"purple") + facet_wrap(vars(cluster))  
#visualizing NumCatalogPurchases  
customers %>% ggplot(aes(NumCatalogPurchases)) + geom_histogram(color =  
"black", fill = "orange") + facet_wrap(vars(cluster))  
#visualizing meat variable  
customers %>% ggplot(aes(meat)) + geom_histogram(color = "black", fill =  
"brown") + facet_wrap(vars(cluster))
```